Table 1: HMB Chronic Perinatal Litter Data Analysis in Female Harlan Sprague Dawley Rats: PND1 Data

Parameter	Control	1000 ppm	3000 ppm	10000 ppm	Trenda
Litter Size <sup>b,d</sup>					
Total	12.55 ± 0.35 [40]	12.65 ± 0.41 [34]	11.63 ± 0.65 [32]	12.53 ± 0.41 [36]	0.716-
Live Male	$6.45 \pm 0.35$ [40]	$6.29 \pm 0.36 [34]$	$5.41 \pm 0.40 [32]$	$6.39 \pm 0.38 [36]$	0.719-
Live Female	$6.00 \pm 0.32$ [40]	$6.21 \pm 0.36 [34]$	$6.03 \pm 0.45 [32]$	$5.94 \pm 0.36 [36]$	0.982+
Live Total	12.45 ± 0.36 [40]	12.50 ± 0.41 [34]	11.44 ± 0.66 [32]	$12.33 \pm 0.43 [36]$	0.610-
% Male per Litter	51.23 ± 2.29 [40]	50.20 ± 2.34 [34]	45.84 ± 2.89 [32]	51.66 ± 2.54 [36]	0.952-
% Male in Dose <sup>e,f</sup>	52 [498]	50 [425]	47 [366]	52 [444]	•
Pup Body Weights <sup>c,d,g</sup>					
Male	$7.32 \pm 0.07 [40]$	$7.38 \pm 0.07 [34]$	$7.22 \pm 0.09 [31]$	7.11 ± 0.08 [35]	0.054-
Female	$6.95 \pm 0.07 [40]$	$7.00 \pm 0.08 [34]$	$6.86 \pm 0.09 [32]$	$6.80 \pm 0.07 [36]$	0.110-
Male+Female	$7.13 \pm 0.07$ [40]	$7.20 \pm 0.07 [34]$	7.01 ± 0.08 [32]	$6.95 \pm 0.07 [36]$	0.044-
Adjusted Pup Body Weights <sup>c,d,h</sup>					
Male	$7.33 \pm 0.07 [40]$	$7.39 \pm 0.07 [34]$	$7.19 \pm 0.08 [31]$	7.11 ± 0.08 [35]	0.017-
Female	$6.96 \pm 0.07 [40]$	$7.02 \pm 0.08 [34]$	$6.82 \pm 0.09 [32]$	$6.81 \pm 0.07 [36]$	0.030-
Male+Female	$7.14 \pm 0.06 [40]$	$7.21 \pm 0.07 [34]$	$6.98 \pm 0.08$ [32]	$6.96 \pm 0.07 [36]$	0.012-

a: P-value and direction of trend

b: Each dose was compared to the control with Shirleys test when a trend was present [P<.01 from Jonckheere's trend test] or with Dunns test when no trend was present [\* = P<0.05, \*\* = P<0.01]

c: Each dose was compared to the control with Williams' test when a trend was present [P < .01 from Jonckheere's trend test] or with Dunnett's test when no trend was present [\* = P < 0.05, \*\* = P < 0.01]

d: Mean ± standard error [number of dams]

e: 100 x [Number of live males in dose group]/[number of live males and females in dose group]
Only litters with at least one live pup included.

f: No statistics done on this endpoint

g: Total pup weight at PND1 divided by number of live pups at PND1

h: Total pup weight at PND1 divided by number of live pups at PND1 then adjusted for total number of live pups at PND1